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Sequence Listing was accepted.

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Reviewer: Anne Corrigan

Timestamp: [year=2008; month=2; day=12; hr=13; min=35; sec=34; ms=766;]

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Application No: 10590441

Version No: 1.0

Input Set:**Output Set:****Started:** 2008-02-12 11:18:30.460**Finished:** 2008-02-12 11:18:32.208**Elapsed:** 0 hr(s) 0 min(s) 1 sec(s) 748 ms**Total Warnings:** 26**Total Errors:** 0**No. of SeqIDs Defined:** 30**Actual SeqID Count:** 30

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Input Set:

Output Set:

Started: 2008-02-12 11:18:30.460
Finished: 2008-02-12 11:18:32.208
Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 748 ms
Total Warnings: 26
Total Errors: 0
No. of SeqIDs Defined: 30
Actual SeqID Count: 30

Error code

Error Description

This error has occurred more than 20 times, will not be displayed

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<120> Highly efficient gene targeting

<130> RJ007P41

<140> 10590441

<141> 2008-02-12

<160> 30

<170> PatentIn version 3.1

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<212> PRT
<213> Homo sapiens

<400> 2

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35 40 45

Glu Asn Lys Asp Glu Ile Ala Leu Val Leu Phe Gly Thr Asp Gly Thr
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Asp Asn Pro Leu Ser Gly Gly Asp Gln Tyr Gln Asn Ile Thr Val His
65 70 75 80

Arg His Leu Met Leu Pro Asp Phe Asp Leu Leu Glu Asp Ile Glu Ser
85 90 95

Lys Ile Gln Pro Gly Ser Gln Gln Ala Asp Phe Leu Asp Ala Leu Ile
100 105 110

Val Ser Met Asp Val Ile Gln His Glu Thr Ile Gly Lys Lys Phe Glu
115 120 125

Lys Arg His Ile Glu Ile Phe Thr Asp Leu Ser Ser Arg Phe Ser Lys
130 135 140

Ser Gln Leu Asp Ile Ile Ile His Ser Leu Lys Lys Cys Asp Ile Ser
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Leu Gln Phe Phe Leu Pro Phe Ser Leu Gly Lys Glu Asp Gly Ser Gly
165 170 175

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Glu	Arg	His	Ser	Ile	His	Trp	Pro	Cys	Arg	Leu	Thr	Ile	Gly	Ser	Asn	
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Leu	Ser	Ile	Arg	Ile	Ala	Ala	Tyr	Lys	Ser	Ile	Leu	Gln	Glu	Arg	Val	
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Lys	Lys	Thr	Trp	Thr	Val	Val	Asp	Ala	Lys	Thr	Leu	Lys	Lys	Glu	Asp	
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Ile	Gln	Lys	Glu	Thr	Val	Tyr	Cys	Leu	Asn	Asp	Asp	Asp	Glu	Thr	Glu	
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Val	Pro	Phe	Ser	Lys	Val	Asp	Glu	Glu	Gln	Met	Lys	Tyr	Lys	Ser	Glu	
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Glu Cys Leu Val Tyr Val Gln Leu Pro Phe Met Glu Asp Leu Arg Gln
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Tyr Met Phe Ser Ser Leu Lys Asn Ser Lys Lys Tyr Ala Pro Thr Glu
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Ala Gln Leu Asn Ala Val Asp Ala Leu Ile Asp Ser Met Ser Leu Ala
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Lys Lys Asp Glu Lys Thr Asp Thr Leu Glu Asp Leu Phe Pro Thr Thr
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Lys Ile Pro Asn Pro Arg Phe Gln Arg Leu Phe Gln Cys Leu Leu His
485 490 495

Arg Ala Leu His Pro Arg Glu Pro Leu Pro Pro Ile Gln Gln His Ile
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Trp Asn Met Leu Asn Pro Pro Ala Glu Val Thr Thr Lys Ser Gln Ile
515 520 525

Pro Leu Ser Lys Ile Lys Thr Leu Phe Pro Leu Ile Glu Ala Lys Lys
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Lys Asp Gln Val Thr Ala Gln Glu Ile Phe Gln Asp Asn His Glu Asp
545 550 555 560

Gly Pro Thr Ala Lys Lys Leu Lys Thr Glu Gln Gly Gly Ala His Phe
565 570 575

Ser Val Ser Ser Leu Ala Glu Gly Ser Val Thr Ser Val Gly Ser Val
580 585 590

Asn Pro Ala Glu Asn Phe Arg Val Leu Val Lys Gln Lys Lys Ala Ser
595 600 605

Phe Glu Glu Ala Ser Asn Gln Leu Ile Asn His Ile Glu Gln Phe Leu
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Asp Thr Asn Glu Thr Pro Tyr Phe Met Lys Ser Ile Asp Cys Ile Arg

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His Phe Trp Glu Ile Val Val Gln Asp Gly Ile Thr Leu Ile Thr Lys						
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Thr Cys Ala Tyr Gln Ile Met Gln Gln Arg Ile Ile Ser Gln Pro Lys						
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Asp Met Met Gly Val Leu Leu Phe Gly Thr Glu Lys Ser Lys Phe Arg						
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Asn	Asp	Asp	Pro	His	Ala	Gly	Asp	Lys	Gln	Ala	Lys	Ser	Ser	Ala	Ala	
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Glu	Gly	Glu	Glu	Gln	Ala	Gln	Val	Val	Gln	Ser	Glu	Thr	Ala	Gln	Val	
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Asp	Phe	Ala	Ala	Arg	Thr	Val	Glu	Lys	Ser	Glu	Leu	Arg	Lys	Gly	Tyr		
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Phe	Pro	Ser	Glu	Glu	Gln	Tyr	Val	Gly	Ser	Thr	Arg	Val	Phe	Ser	Ala		
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Val	Ala	Arg	Glu	Asn	Ala	His	Pro	Val	Met	Val	Ala	Ile	Phe	Pro	Ser		
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His	Val	Thr	Ala	Pro	Pro	Arg	Pro	Ala	Asp	Glu	Leu	Thr	Asp	Gln	Met		
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Arg	Lys	Tyr	Pro	Asn	Pro	Ser	Leu	Gln	Trp	His	Tyr	Lys	Ile	Leu	Gln		
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Ala	Lys	Ala	Leu	Asp	Glu	Glu	Thr	Pro	Asp	Ala	Met	Asp	Asp	Val	Thr		
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Arg Ala Lys Pro Ser Lys Lys Ala Ala Ser Gly Gly Gly Gly Pro Ala
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Asn Ser Asn Ala Asp Leu Lys Lys Ala Phe Glu Gln Gly Thr Leu Gly
595 600 605

Lys Met Thr Val Ala Glu Leu Lys Asp Ile Met Ala Ser Lys Gly Ile
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